

SCORE Search Results Details for Application  
10668767 and Search Result us-10-668-767-  
128\_copy\_1\_1700.p2n.rnpbm.

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## Retrieve Application List

## SCORE System Overview

SCORE  
FAQ

Comments /  
Suggestions

This page gives you Search Results detail for the Application 10668767 and Search Result us-10-668-767-128\_copy\_1\_1700.p2n.rnpbm.

start

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GenCore version 5.1.9  
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OM protein - nucleic search, using frame\_plus\_p2n model

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Run on:      October 23, 2006, 08:16:57 ; Search time 3663 Seconds
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## SUMMARIES

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1	8985	100.0	15387	8	US-10-668-767-127	Sequence 127, App
2	8970	99.8	15405	8	US-10-668-767-129	Sequence 129, App
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4	8872.5	98.7	15679	8	US-10-668-767-1	Sequence 1, Appli
5	8734	97.2	15303	8	US-10-668-767-145	Sequence 145, App
6	7112	79.2	15413	8	US-10-668-767-7	Sequence 7, Appli
7	6964.5	77.5	16363	8	US-10-668-767-9	Sequence 9, Appli
8	6920.5	77.0	15315	8	US-10-668-767-5	Sequence 5, Appli
9	6887	76.6	15606	13	US-11-097-143-22562	Sequence 22562, A
10	6841.5	76.1	15845	8	US-10-668-767-3	Sequence 3, Appli
c 11	6196.5	69.0	24236	13	US-11-097-143-22561	Sequence 22561, A
12	3817	42.5	14302	8	US-10-276-774-552	Sequence 552, App
13	3817	42.5	15731	10	US-10-887-553A-490	Sequence 490, App
14	3660	40.7	15359	8	US-10-276-774-500	Sequence 500, App
15	3660	40.7	15359	10	US-10-887-553A-489	Sequence 489, App
16	3660	40.7	15359	10	US-10-450-763-4960	Sequence 4960, Ap
17	3655	40.7	15563	8	US-10-764-425-48	Sequence 48, Appl
18	3655	40.7	15563	10	US-10-887-553A-491	Sequence 491, App
19	3653.5	40.7	14620	15	US-11-044-111-17	Sequence 17, Appl
20	3640	40.5	15564	15	US-11-000-688-1500	Sequence 1500, Ap
21	3606	40.1	15583	10	US-10-450-763-12305	Sequence 12305, A
22	3606	40.1	15820	10	US-10-450-763-23337	Sequence 23337, A
23	3536	39.4	15453	9	US-10-723-860-5796	Sequence 5796, Ap
24	3525	39.2	4078	7	US-10-305-720-1154	Sequence 1154, Ap
25	2709.5	30.2	3168	15	US-11-044-111-8	Sequence 8, Appli
26	698.5	7.8	1800	10	US-10-450-763-23334	Sequence 23334, A
27	603	6.7	717	15	US-11-044-111-18	Sequence 18, Appl
28	576.5	6.4	636	15	US-11-044-111-19	Sequence 19, Appl
29	424	4.7	524	15	US-11-044-111-20	Sequence 20, Appl
30	327	3.6	10129	15	US-11-044-111-21	Sequence 21, Appl
31	324.5	3.6	597	7	US-10-029-386-2405	Sequence 2405, Ap
32	324	3.6	11806	8	US-10-367-094-123	Sequence 123, App
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37	319.5	3.6	10524	8	US-10-240-425-1212	Sequence 1212, Ap
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c 40	318.5	3.5	535	6	US-10-027-632-59310	Sequence 59310, A
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c 43	318.5	3.5	643	7	US-10-027-632-51124	Sequence 51124, A
44	299	3.3	8806	8	US-10-152-319A-1612	Sequence 1612, Ap
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## ALIGNMENTS

## RESULT 1

US-10-668-767-127

; Sequence 127, Application US/10668767

; Publication No. US20040171114A1

; GENERAL INFORMATION:

; APPLICANT: Caspar, Timothy

; APPLICANT: Cordova, Daniel

; APPLICANT: Gutteridge, Steven

; APPLICANT: Rauh, James

; APPLICANT: Smith, Rejane

; APPLICANT: Tao, Yong

; APPLICANT: Wu, Lihong

; TITLE OF INVENTION: Isolation and Uses of Ryanodine Receptors

; FILE REFERENCE: BB1533 US NA

; CURRENT APPLICATION NUMBER: US/10/668,767

; CURRENT FILING DATE: 2003-09-23

; PRIOR APPLICATION NUMBER: 60/412,795

; PRIOR FILING DATE: 2002-09-23

; PRIOR APPLICATION NUMBER: 60/427,324

; PRIOR FILING DATE: 2002-11-18

; NUMBER OF SEQ ID NOS: 149

; SOFTWARE: PatentIn version 3.1

; SEQ ID NO 127

; LENGTH: 15387

; TYPE: DNA

; ORGANISM: Heliothis virescens

; FEATURE:

; NAME/KEY: CDS

; LOCATION: (1)..(15384)

; OTHER INFORMATION:

; FEATURE:

; OTHER INFORMATION: pXL-Hv7

US-10-668-767-127

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## SUMMARIES

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7	279.5	3.1	8404	8	US-11-266-748A-28705	Sequence 28705, A
8	235.5	2.6	1406	8	US-11-266-748A-31954	Sequence 31954, A
9	193.5	2.2	2084	8	US-11-266-748A-31119	Sequence 31119, A
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12	183.5	2.0	4226	8	US-11-293-697-1236	Sequence 1236, Ap
13	180.5	2.0	1450	8	US-11-266-748A-186649	Sequence 186649, A
14	180.5	2.0	1937	1	US-09-949-925-61	Sequence 61, Appl
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19	155	1.7	1991	8	US-11-266-748A-251793	Sequence 251793, A
c 20	155	1.7	1991	8	US-11-266-748A-312310	Sequence 312310, A
c 21	154.5	1.7	583	8	US-11-266-748A-377158	Sequence 377158, A
22	154.5	1.7	583	8	US-11-266-748A-460537	Sequence 460537, A
23	154	1.7	2667	8	US-11-266-748A-23535	Sequence 23535, A
24	149	1.7	2947	8	US-11-266-748A-26565	Sequence 26565, A
25	144.5	1.6	349	8	US-11-266-748A-373619	Sequence 373619, A
c 26	144.5	1.6	349	8	US-11-266-748A-456998	Sequence 456998, A
27	143	1.6	813	8	US-11-266-748A-193768	Sequence 193768, A
28	143	1.6	813	8	US-11-266-748A-227276	Sequence 227276, A
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## ALIGNMENTS

## RESULT 1

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; Sequence 57467, Application US/11266748A

; Publication No. US20060134663A1

; GENERAL INFORMATION:

; APPLICANT: Harkin, Paul

; APPLICANT: Johnston, Patrick

; APPLICANT: Mulligan, Karl

; TITLE OF INVENTION: Transcriptome Microarray Technology and

; TITLE OF INVENTION: Methods of Using the Same

; FILE REFERENCE: 55815-0102 (319189)

; CURRENT APPLICATION NUMBER: US/11/266,748A

; CURRENT FILING DATE: 2005-11-03

; PRIOR APPLICATION NUMBER: EP 04105479.2

; PRIOR FILING DATE: 2004-11-03

; PRIOR APPLICATION NUMBER: EP 04105482.6

; PRIOR FILING DATE: 2004-11-03

; PRIOR APPLICATION NUMBER: EP 04105483.4

; PRIOR FILING DATE: 2004-11-03

; PRIOR APPLICATION NUMBER: EP 04105507.0

; PRIOR FILING DATE: 2004-11-03

; PRIOR APPLICATION NUMBER: EP 04105485.9

; PRIOR FILING DATE: 2004-11-03

; PRIOR APPLICATION NUMBER: EP 04105484.2

; PRIOR FILING DATE: 2004-11-03

; PRIOR APPLICATION NUMBER: US 60/662,276

; PRIOR FILING DATE: 2005-03-14

; PRIOR APPLICATION NUMBER: US 60/700,293

; PRIOR FILING DATE: 2005-07-18

; NUMBER OF SEQ ID NOS: 483996

; SOFTWARE: PatentIn version 3.3

; SEQ ID NO 57467

; LENGTH: 15563

; TYPE: DNA

; ORGANISM: Homo Sapiens

US-11-266-748A-57467

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44	132.5	1.5	2655	2	US-08-470-566B-17	Sequence 17, Appl
45	132.5	1.5	2655	2	US-08-470-566B-26	Sequence 26, Appl

## ALIGNMENTS

## RESULT 1

US-08-785-420-1

; Sequence 1, Application US/08785420

; Patent No. 6001976

; GENERAL INFORMATION:

; APPLICANT: MacLennan, David H

; APPLICANT: O'Brien, Peter J.

; TITLE OF INVENTION: DIAGNOSIS FOR PORCINE MALIGNANT

; TITLE OF INVENTION: HYPERTHERMIA

; NUMBER OF SEQUENCES: 3

; CORRESPONDENCE ADDRESS:

; ADDRESSEE: Bell, Seltzer, Park &amp; Gibson

; STREET: P.O. Drawer 34009

; CITY: Charlotte,

; STATE: No. 6001976th Carolina 28234

; COUNTRY: U.S.A.

; ZIP: 28234

; COMPUTER READABLE FORM:

; MEDIUM TYPE: Floppy disk

; COMPUTER: IBM PC compatible

; OPERATING SYSTEM: PC-DOS/MS-DOS

; SOFTWARE: PatentIn Release #1.0, Version #1.25

; CURRENT APPLICATION DATA:

; APPLICATION NUMBER: US/08/785,420

; FILING DATE:

; CLASSIFICATION:

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER: 08/249,388

; FILING DATE:

; APPLICATION NUMBER: US 08/030,159

; FILING DATE: 15-MAR-1993

; ATTORNEY/AGENT INFORMATION:

; NAME: Layton, Jr., Samuel G.

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; REFERENCE/DOCKET NUMBER: 3477-73

; TELECOMMUNICATION INFORMATION:

; TELEPHONE: 704-377-1561

; TELEFAX: 704-334-2014

; TELEX: 57-5102

; INFORMATION FOR SEQ ID NO: 1:

; SEQUENCE CHARACTERISTICS:

; LENGTH: 15378 base pairs

; TYPE: nucleic acid

; STRANDEDNESS: single

; TOPOLOGY: linear

; MOLECULE TYPE: DNA (genomic)

; HYPOTHETICAL: NO

; ANTI-SENSE: NO

; IMMEDIATE SOURCE:

; CLONE: Porcine RYR1 Gene

; POSITION IN GENOME:

; UNITS: bp

US-08-785-420-1